1646

RAW SEQUENCE LISTING   DATE: 11/13/2000
PATENT APPLICATION: US/09/509,196A   TIME: 13:25:59

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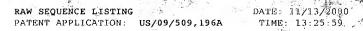Input Set : A:\1871-121.app
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 3  <110> APPLICANT: DALY, Roger J.
 4        SUTHERLAND, Robert L.
 6  <120> TITLE OF INVENTION: A Potential Effector for the Grb7 Family of Signalling
 7        Proteins
 9  <130> FILE REFERENCE: 1871-129
11  <140> CURRENT APPLICATION NUMBER: 09/509,196A
12  <141> CURRENT FILING DATE: 2000-03-23
14  <150> PRIOR APPLICATION NUMBER: P09388
15  <151> PRIOR FILING DATE: 1997-09-23
17  <150> PRIOR APPLICATION NUMBER: PCT AU98/00795
18  <151> PRIOR FILING DATE: 1998-09-23
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41  gctatagact tggctcccac accacagtta aaagaaagat tagcatatga atttaaaggc 720
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62 aagtataatg catctctcaa tgccacggac aaatgggctt tcacaccttt gcacgaagca 1980
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99                 20                  25                  30
101 Tyr Thr Pro Leu His Glu Ala Ala Ile Lys Gly Lys Ile Asp Val Cys
102                35                  40                  45
104 Ile Val Leu Leu Gln His Gly Ala Glu Pro Thr Ile Arg Asn Thr Asp
105       50                  55                  60
107 Gly Arg Thr Ala Leu Asp Leu Ala Asp-Pro Ser Ala Lys Ala Val Leu
108   65                  70                  75                  80
110 Thr Gly Glu Tyr Lys Lys Asp Glu Leu Leu Glu Ser Ala Arg Ser Gly
111                85                  90                  95
113 Asn Glu Glu Lys Met Met Ala Leu Leu Thr Pro Leu Asn Val Asn Cys
114                100                 105                 110
116 His Ala Ser Asp Gly Arg Lys Ser Thr Pro Leu His Leu Ala Ala Gly
117            115                 120                 125
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119 Tyr Asn Arg Val Lys Ile Val Gln Leu Leu Leu Gln His Gly Arg Asp
120     130             135               140
122 Val His Ala Lys Asp Lys Gly Asp Leu Val Pro Leu His Asn Ala Cys
123 145             150             155               160
125 Ser Tyr Gly His Tyr Glu Val Thr Glu Leu Leu Val Lys His Gly Gly
126             165             170               175
128 Cys Val Asn Ala Met Asp Leu Trp Gln Phe Thr Pro Leu His Glu Ala
129             180             185               190
131 Ala Ser Lys Asn Arg Val Glu Val Cys Ser Leu Leu Leu Ser Tyr Gly
132     195             200                 205
134 Ala Asp Pro Thr Leu Leu Asn Cys Lys Asn Lys Ser Ala Ile Asp Leu
135     210             215                 220
137 Ala Pro Thr Pro Gln Leu Lys Glu Arg Leu Ala Tyr Glu Phe Lys Gly
138 225             230             235               240
140 His Ser Leu Leu Gln Ala Ala Arg Glu Ala Asp Val Thr Arg Ile Lys
141             245             250               255
143 Lys His Leu Ser Leu Glu Met Val Asn Phe Lys His Pro Gln Thr His
144             260             265             270
146 Glu Thr Ala Leu His Cys Ala Ala Ala Ser Pro Tyr Pro Lys Arg Lys
147     275             280             285
149 Gln Ile Cys Glu Leu Leu Leu Arg Lys Gly Ala Asn Ile Asn Glu Lys
150     290             295               300
152 Thr Lys Glu Phe Leu Thr Pro Leu His Val Ala Ser Glu Lys Ala His
153 305             310             315               320
155 Asn Asp Val Val Glu Val Val Val Lys His Glu Ala Lys Val Asn Ala
156             325             330               335
158 Leu Asp Asn Leu Gly Gln Thr Ser Leu His Arg Ala Ala Tyr Cys Gly
159             340             345             350
161 His Leu Gln Thr Cys Arg Leu Leu Leu Ser Tyr Gly Cys Asp Pro Asn
162     355             360             365
164 Ile Ile Ser Leu Gln Gly Phe Thr Ala Leu Gln Met Gly Asn Glu Asn
165     370             375             380
167 Val Gln Gln Leu Leu Gln Glu Gly Ile Ser Leu Gly Asn Ser Glu Ala
168 385             390             395               400
170 Asp Arg Gln Leu Leu Glu Ala Ala Lys Ala Gly Asp Val Glu Thr Val
171             405             410             415
173 Lys Lys Leu Cys Thr Val Gln Ser Val Asn Cys Arg Asp Ile Glu Gly
174             420             425             430
176 Arg Gln Ser Thr Pro Leu His Phe Ala Ala Gly Tyr Asn Arg Val Ser
177     435             440             445
179 Val Val Glu Tyr Leu Leu Gln His Gly Ala Asp Val His Ala Lys Asp
180     450             455             460
182 Lys Gly Gly Leu Val Pro Leu His Asn Ala Cys Ser Tyr Gly His Tyr
183 465             470             475               480
185 Glu Val Ala Glu Leu Leu Val Lys His Gly Ala Val Val Asn Val Ala
186             485             490               495
188 Asp Leu Trp Lys Phe Thr Pro Leu His Glu Ala Ala Ala Lys Gly Lys
189             500             505             510
191 Tyr Glu Ile Cys Lys Leu Leu Leu Gln His Gly Ala Asp Pro Thr Lys
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197 Thr Asp Ile Gln Asp Leu Leu Arg Gly Asp Ala Ala Leu Leu Asp Ala
198 545                   550                   555                   560
200 Ala Lys Lys Gly Cys Leu Ala Arg Val Lys Lys Leu Ser Ser Pro Asp
201             565                   570                   575
203 Asn Val Asn Cys Arg Asp Thr Gln Gly Arg His Ser Thr Pro Leu His
204             580                   585                   590
206 Leu Ala Ala Gly Tyr Asn Asn Leu Glu Val Ala Glu Tyr Leu Leu Gln
207         595                   600                   605
209 His Gly Ala Asp Val Asn Ala Gln Asp Lys Gly Gly Leu Ile Pro Leu
210         610                   615                   620
212 His Asn Ala Ala Ser Tyr Gly His Val Asp Val Ala Ala Leu Leu Ile
213 625                   630                   635                   640
215 Lys Tyr Asn Ala Ser Leu Asn Ala Thr Asp Lys Trp Ala Phe Thr Pro
216             645                   650                   655
218 Leu His Glu Ala Ala Gln Lys Gly Arg Thr Gln Leu Cys Ala Leu Leu
219             660                   665                   670
221 Leu Ala His Gly Ala Asp Pro Thr Leu Lys Asn Gln Glu Gly Gln Thr
222         675                   680                   685
224 Pro Leu Asp Leu Val Ser Ala Asp Asp Val Ser Ala Leu Leu Thr Ala
225         690                   695                   700
227 Ala Met Pro Pro Ser Ala Leu Pro Ser Cys Tyr Lys Pro Gln Val Leu
228 705                   710             .   715                   720
230 Asn Gly Val Arg Ser Pro Gly Ala Thr Ala Asp Ala Leu Ser Ser Gly
231             725                   730                   735
233 Pro Ser Ser Pro Ser Ser Leu Ser Ala Ala Ser Ser Leu Asp Asn Leu
234             740                   745                   750
236 Ser Gly Ser Phe Ser Glu Leu Ser Ser Val Val Ser Ser Ser Gly Thr
237         755                   760                   765
239 Glu Gly Ala Ser Ser Leu Glu Lys Lys Glu Val Pro Gly Val Asp Phe
240         770                   775                   780
242 Ser Ile Thr Gln Phe Val Arg Asn Leu Gly Leu Glu His Leu Met Asp
243 785                   790                   795                   800
245 Ile Phe Glu Arg Glu Gln Ile Thr Leu Asp Val Leu Val Glu Met Gly
246             805                   810                   815
248 His Lys Glu Leu Lys Glu Ile Gly Ile Asn Ala Tyr Gly His Arg His
249             820                   825                   830
251 Lys Leu Ile Lys Gly Val Glu Arg Leu Ile Ser Gly Gln Gln Gly Leu
252         835                   840                   845
254 Asn Pro Tyr Leu Thr Leu Asn Thr Ser Gly Ser Gly Thr Ile Leu Ile
255 850                   855                   860
257 Asp Leu Ser Pro Asp Asp Lys Glu Phe Gln Ser Val Glu Glu Glu Met
258 865                   870                   875                   880
260 Gln Ser Thr Val Arg Glu His Arg Asp Gly Gly His Ala Gly Gly Ile
261             885                   890                   895
263 Phe Asn Arg Tyr Asn Ile Leu Lys Ile Gln Lys Val Cys Asn Lys Lys
264             900                   905                   910
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266 Leu Trp Glu Arg Tyr Thr His Arg Arg Lys Glu Val Ser Glu Glu Asn
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269 His Asn His Ala Asn Glu Arg Met Leu Phe His Gly Ser Pro Phe Val
270         930                 935                 940
272 Asn Ala Ile Ile His Lys Gly Phe Asp Glu Arg His Ala Tyr Ile Gly
273 945                 950                 955                 960
275 Gly Met Phe Gly Ala Gly Ile Tyr Phe Ala Glu Asn Ser Ser Lys Ser
276             965                 970                 975
278 Asn Gln Tyr Val Tyr Gly Ile Gly Gly Gly Thr Gly Cys Pro Val His
279             980                 985                 990
281 Lys Asp Arg Ser Cys Tyr Ile Cys His Arg Gln Leu Leu Phe Cys Arg
282         995                 1000                1005
284 Val Thr Leu Gly Lys Ser Phe Leu Gln Phe Ser Ala Met Lys Met Ala
285     1010                1015                1020
287 His Ser Pro Pro Gly His His Ser Val Thr Gly Arg Pro Ser Val Asn
288 1025                1030                1035                1040
290 Gly Leu Ala Leu Ala Glu Tyr Val Ile Tyr Arg Gly Glu Glu Ala Tyr
291             1045                1050                1055
293 Pro Glu Tyr Leu Ile Thr Tyr Gln Ile Met Arg Pro Glu Gly Met Val
294         1060                1065                1070
296 Asp Gly
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VERIFICATION SUMMARY   DATE: 11/13/2000
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